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Fast volumetric imaging of ethanol metabolism in rat liver with hyperpolarized [1-¹³C]-pyruvate

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Abstract

Rapid, volumetric imaging of hyperpolarized ¹³C compounds allows the real time measurement of metabolic activity and can be useful in distinguishing between normal and diseased tissues. This work extends a fast 2D under-sampled spiral magnetic resonance spectroscopic imaging (MRSI) sequence to provide volumetric coverage, acquiring a 16×16×12 matrix with a nominal 5 mm isotropic resolution in 4.5 s. The rapid acquisition enables a high temporal resolution for dynamic imaging. This dynamic 3D MRSI method was used to investigate hyperpolarized [1-¹³C]-pyruvate metabolism modulated by the administration of ethanol in rat liver. A significant increase in the pyruvate to lactate conversion was observed in the liver due to the greater availability of NADH from ethanol metabolism.

Keywords

hyperpolarized ¹³C; dynamic metabolic imaging; fast volumetric imaging; liver ethanol; spiral MRSI

Introduction

Hyperpolarized ¹³C magnetic resonance spectroscopic imaging (MRSI) is a powerful tool for non-invasive in vivo measurement of metabolism in real time. Several studies have demonstrated metabolic imaging of hyperpolarized [1-¹³C]-pyruvate (Pyr) and its downstream products lactate (Lac) and alanine (Ala) for a variety of applications, including cancer diagnosis, monitoring therapeutic response, and cardiovascular pathologies [1–7]. One of the challenges of imaging hyperpolarized ¹³C labeled compounds is the rapid decay of the hyperpolarized magnetization. The non-equilibrium magnetization decays to thermal polarization with relaxation time constant T₁, which is typically on the order of 20–40 s in vivo. Several fast spectroscopic imaging techniques have been developed to capture the metabolic information within the limited lifetime of the hyperpolarized signal [8–17]. Rapid acquisition methods are especially needed in order to capture the metabolic kinetics with adequate temporal resolution. One such fast sequence is the 2D spectrally under-sampled

spiral MRSI [9–10], which allows imaging of a slice in less than one second. However, 2D multi-slice approaches suffer from the problems of gaps between slices and chemical shift displacement. For example, chemical shift mis-registration between slice locations of Pyr and Lac, which are 390 Hz apart at 3T, using an RF excitation bandwidth of 2.3 kHz is 1.7 mm for a 10-mm thick slice. Though chemical shift displacement is also present for the Pyr and Lac slab profiles in a 3D acquisition, the phase encoding provides contiguous slices without mis-registration at the expense of some potential slice aliasing at the edge of the field of view (FOV). The 3D approach also allows phase modulating in the slice encode direction during reconstruction to shift the slice locations by a fraction of the slice thickness, as well as retrospectively reformatting the image plane.

The conversion of Pyr to Lac takes place via the enzyme lactate dehydrogenase (LDH) and the concomitant oxidation of NADH to nicotinamide adenine dinucleotide (NAD⁺). At a sufficiently high dose of Pyr, this reaction is saturated, with both LDH activity and NADH availability being rate-limiting steps. The metabolism of ethanol in the liver takes place through the breakdown of ethanol to acetaldehyde and acetaldehyde to acetate. These reactions are catalyzed by the enzymes alcohol dehydrogenase and acetaldehyde dehydrogenase respectively, and both of these reactions reduce the coenzyme NAD⁺ to NADH. Thus, ethanol consumption leads to accumulation of NADH in the liver [18–20], and provides a mechanism by which NADH availability can be altered and used to modulate the Pyr-to-Lac metabolic pathway [21].

Spielman et al. [21] have previously reported MRS measurements of hyperpolarized [1-¹³C]-Pyr to study rat liver metabolism modulated by ethanol. Dynamic ¹³C MRS was used to measure the conversion rates of Pyr to Lac and Ala in the absence and presence of ethanol. However, the MRS data were not spatially localized within a slice, which contained vascular and other structures in addition to the targeted organs, and may underestimate the conversion rates due to the large Pyr signal contribution from the vasculature. Dynamic 3D ¹³C MRSI can provide improved rate-constant estimates by allowing the analysis of spectra from organ-specific regions-of-interest (ROIs). This work extends the fast 2D spiral MRSI sequence to provide volumetric coverage and applies it to investigate Pyr metabolism in rat liver modulated by ethanol.

Method

The polarized sample consisted of 25 μ L of a mixture of 14-M [1-¹³C] pyruvic acid and 15-mM Ox063 trityl radical, to which 3 μ L of a 1:50 dilution of Dotarem (Guerbet, France) was added prior to polarization. The sample was polarized via dynamic nuclear polarization [22] using a HyperSense system (Oxford Instruments Molecular Biotoools, Oxford, UK) to achieve approximately 24% liquid-state polarization at dissolution. The polarized sample was dissolved with a solution of 80 mM NaOH mixed with 40 mM Tris buffer and 0.1 g/L EDTA-Na₂, leading to an 80-mM solution of hyperpolarized pyruvate with a pH of 7.4–7.5. Healthy male Wistar rats (n=5, 275–352 g body weight) were injected with 3 mL of the hyperpolarized pyruvate solution through a tail vein catheter at a rate of approximately 0.2 mL/s. All animal procedures were approved by the Institutional Animal Care and Use Committee.

All experiments were performed on a clinical 3T Signa MR scanner (GE Healthcare, Waukesha, WI) equipped with self-shielded gradients (40 mT/m, 150 mT/m/ms). A custom-built dual-tuned (¹H/¹³C) quadrature rat coil (inner diameter=80 mm, length=90 mm), operating at 127.7 MHz and 32.1 MHz, respectively, was used for both RF excitation and signal reception. The transmit ¹³C RF power was calibrated using a reference phantom containing an 8-M solution of ¹³C-urea, which was placed on top of the animal. Single-shot

fast spin-echo ^1H MR images with nominal in-plane resolution of 0.47 mm and 2-mm slice thickness were acquired in the axial, sagittal, and coronal planes throughout the scan session as anatomical references for prescribing the ^{13}C -MRSI experiments. Additional ^1H 3D spoiled gradient echo (SPGR) images (0.625 mm in-plane resolution, 1.25 mm slice thickness, 96 slices) were also acquired matching the ^{13}C MRSI prescription for overlay of the metabolic images.

The 2D under-sampled spiral spectroscopic imaging sequence described in [9–10] was extended for volumetric coverage. A spiral readout trajectory encoded the spectral and two spatial (x–y) dimensions using 32 echoes and 3 spiral interleaves. The slice (z) direction used 12 phase encoding steps in a centric order. Imaging parameters were: FOV=80×80×60 mm³, 5×5×5 mm³ nominal resolution, spectral width=280 Hz, 42-mm slab excitation, 36 excitations/volume, flip angle=5.625°, TE=2.3 ms, readout duration=114.3 ms, nominal spectral resolution=8.7 Hz, total acquisition time=4.5 s. The cumulative effect of 36 excitations of 5.625° each leads to an effective flip angle of 32.8° for the volume acquisition. The effective flip angle was estimated as $\alpha_{\text{eff}} = \cos^{-1}(\cos^N \alpha_k)$, where $\alpha_k = 5.625^\circ$ and $N=36$. This calculation assumes instantaneous successive excitations, ignoring any changes in pyruvate magnetization during the time between excitations. Dynamic ^{13}C MRSI data were acquired from a 3D volume that included both liver and kidney every 5 s for 80 s, starting at the same time as the Pyr injection. The time from dissolution to start of injection was kept constant at 24 s.

After acquiring a baseline ^{13}C MRSI measurement, 1.0 g/kg of a 20% ethanol solution was injected into the tail vein at the rate of 1 mL/min. Another ^{13}C MRSI measurement was acquired approximately 45 min after the ethanol injection to achieve a targeted steady-state blood alcohol level (BAL) of approximately 100 mg/dL at the time of the second acquisition. At the end of the exam, blood was collected for BAL analysis.

The data were reconstructed similarly as the 2D case described in [9–10] with an additional apodization and Fourier transform step in the slab-select direction. The data were zero-filled by a factor of 2 in each spatial direction. Due to spectral undersampling and aliasing of the metabolites, a separate reconstruction is required for each metabolite. With the center frequency approximately at alanine, pyruvate and lactate were aliased once in opposite directions. In each reconstruction, one of the spectral resonances is corrected for the chemical shift phase accrual and reconstructed ‘in-focus’ while the aliased resonances are blurred. A zero-order phase correction is then applied to put the ‘in-focus’ peak into absorption mode. The details of the reconstruction are given in [9–10].

Metabolic maps for Pyr, Lac, and Ala were calculated by integrating the signal within ± 20 Hz around each peak in absorption mode. The metabolic images were corrected for differences in the solid-state polarization level of each sample. ROIs were drawn manually for liver (mean ROI size was 1.22 cm³), kidney (0.94 cm³) and vasculature (0.28 cm³) in both the liver and kidney slices to calculate the mean time-resolved signal intensities for Pyr, Lac, and Ala. The liver and kidney ROIs were also used to calculate signal-to-noise ratios (SNR) in the time-averaged metabolic images. The noise was measured as the standard deviation in a region of the FOV outside the animal in a slice outside the excited slab. Additionally, a large ROI encompassing the entire animal (17.8 cm³) was used to generate whole-slice time-courses for the liver and kidney slices. The metabolite time-courses were then fit using a three-site exchange model to estimate the apparent Pyr-to-Lac (k_{pl}) and Pyr-to-Ala (k_{pa}) rate constants. The model used here was the same as described in [21], except that the measured Pyr time-curve was used as the bolus input function into the model. It used linear gray-box modeling (idgrey, MatLab, MathWorks, Natick, MA) to estimate the parameters by minimizing the sum of squares of the residual error at each time point. The

parameters fit by this model were apparent k_{pl} and k_{pa} exchange rate constants, and apparent T_1 decay time constants for Pyr (T_{1pyr}), Lac (T_{1lac}) and Ala (T_{1ala}). In order to simplify the model, the backward exchange rates, k_{lp} and k_{ap} , were set to zero. The apparent exchange rate constants include contributions from metabolic flux, isotopic exchange, and transport kinetics. Similarly, the apparent T_1 decay constant represents polarization loss over time from T_1 relaxation as well as from RF excitation and inflow/outflow of spins.

A statistical analysis of the estimated metabolic rate constants pre- and post-ethanol was performed to identify significant changes. Due to the small sample size and non-normal distribution, a non-parametric Wilcoxon signed rank test was used for hypothesis testing.

Results and Discussion

Figure 1 shows representative post-ethanol ^{13}C metabolic maps of Pyr, Lac, and Ala from the 3D volume covering kidney and liver superimposed onto corresponding ^1H 3D SPGR slices. The central 12 of the 24 reconstructed slices are shown here. The ^{13}C images were averaged over all of the 16 time-points acquired during the 80-s scan. The time series of ^{13}C -Lac maps from the kidney and liver slices shown in Fig. 2 illustrate the temporal dynamics and spatial distribution of this metabolite. The fast acquisition achieved by spiral MRSI enabled a temporal resolution of 5 s for dynamic imaging. These figures present an example of time-resolved volumetric imaging, which allows the simultaneous characterization of metabolic profiles from different tissues.

The baseline (pre-ethanol) SNR values were calculated for the time-averaged Pyr, Lac and Ala in liver and kidney. For liver, the SNR of Pyr, Lac, and Ala were 63.8 ± 10.7 (mean \pm standard deviation, $n=5$), 37.5 ± 6.6 , and 29.3 ± 3.2 respectively. For kidney, the SNR of Pyr, Lac, and Ala were 304.7 ± 79.1 , 70.4 ± 8.6 , and 30.8 ± 3.4 respectively. The kidney SNR was measured in the right kidney, which lies relatively close to the aorta and vena cava. The pyruvate level in vasculature is much higher than that in the kidney, and at the 5 mm spatial resolution used here, there can be some pyruvate signal contribution from the vasculature into the kidney ROI. This may be the cause for the high inter-subject variability seen in the pyruvate SNR in kidney. This signal contamination from vasculature is much lower for lactate and alanine, and hence could also lead to underestimation of apparent k_{pl} and k_{pa} . Imaging with higher spatial resolution would help alleviate this problem.

Figure 3 shows the time-averaged lactate spectra, normalized to pyruvate, pre- and post-ethanol in liver and in kidney. The spectra from the liver show higher lactate signal post-ethanol compared to pre-ethanol, as expected from the NADH accumulation due to ethanol metabolism. The alanine signal was unchanged. The kidney lactate signal pre- and post-ethanol was also similar. The spectra shown here are from the lactate reconstruction where lactate is corrected for the chemical shift phase accrual and reconstructed 'in-focus' while the aliased resonances are blurred

Figure 4 plots the time-courses of the mean Pyr, Lac, and Ala signals from ROIs in the liver and the kidney, measured pre- and post-ethanol, from one animal. The corresponding fits with the apparent rate constants are also shown. The liver plots clearly show increased lactate signal post-ethanol compared to pre-ethanol, while there was little change in the kidney lactate signal. Alanine signal also was similar before and after ethanol in both organs. The estimated apparent k_{pl} and k_{pa} rate constants for all 5 rats are summarized in Fig. 5 and in Table 1. The BALs for the 5 animals ranged from 81.1 to 101.6 mg/dL. The average apparent liver k_{pl} increased from $0.049 \pm 0.007 \text{ s}^{-1}$ (mean \pm standard deviation) pre-ethanol to $0.064 \pm 0.012 \text{ s}^{-1}$ post-ethanol. This increase in k_{pl} was found to be statistically significant ($P < 0.044$), and is consistent with the hypothesis that NADH levels are rate limiting for liver Pyr-to-Lac conversion at the given Pyr dose [21]. As a control, no

significant changes were observed for the labeled lactate production in kidney or vasculature ROIs or for alanine production in any of the tissues. The inter-subject variability leads to high standard deviations in Table 1, but the intra-subject comparison between pre- and post-ethanol in Fig. 5 clearly shows the change in rate constants.

Compared to the slice-selective MRS measurements in [21], the organ specific apparent k_{pl} and k_{pa} values obtained here at baseline were considerably higher in the liver but similar in the kidney. For example, the average apparent k_{pl} at baseline in liver was $0.049 \pm 0.007 \text{ s}^{-1}$ for the 3D MRSI and $0.014 \pm 0.003 \text{ s}^{-1}$ for the slice-selective MRS in ref. 21. The corresponding values in kidney were $0.018 \pm 0.004 \text{ s}^{-1}$ for the 3D MRSI and $0.012 \pm 0.002 \text{ s}^{-1}$ for the slice-selective MRS. The greater difference in liver k_{pl} is presumably because the fraction of Pyr signal contribution from the vasculature to the total Pyr signal in the slice is higher for the liver slice than for the kidney slice. This effect is also seen in the Pyr images in Fig. 1. The Pyr signal in the liver is much lower compared to that in the blood vessel in the same slice, and is below the threshold used for display of the ^{13}C maps. However, the Pyr signal in the kidney relative to the vessel in the kidney slice is easily visible. Due to the large contribution of the vascular Pyr signal, the whole-slice ROI is a better comparison to the slice-selective estimates. For the whole-slice time-courses generated from the 3D MRSI for liver and kidney slices, the apparent k_{pl} were $0.015 \pm 0.003 \text{ s}^{-1}$ for liver and $0.016 \pm 0.006 \text{ s}^{-1}$ for kidney, and the apparent k_{pa} were $0.010 \pm 0.002 \text{ s}^{-1}$ for liver and $0.011 \pm 0.003 \text{ s}^{-1}$ for kidney, all of which are similar to the slice-selective MRS estimates from [18]. These values demonstrate how the organ-specific rate constants are underestimated by slice-selective dynamic data due to the large vascular Pyr component.

The three-site exchange model used for fitting does not account for the RF excitation scheme used for data acquisition. The slice-selective measurements in [21] used a single RF excitation with a 5° flip angle for each time-point, while the 3D imaging here used 36 excitations of 5.6° each, resulting in a much higher effective flip angle per time-point. Also, the 3D sequence had an acquisition duration of 4.5 s and a TR of 5 s with inflow occurring throughout, as compared to the short 409.6 ms readout and TR of 3 s in [21]. These differences in the RF sampling schemes and the inflow effects could lead to differences in the absolute values of the apparent rate constant and T_1 decay estimates. However, the impact of the RF sampling is the same for both pre- and post-ethanol measurements, and should have only a small effect on the change in apparent k_{pl} and k_{pa} from pre- to post-ethanol infusion.

A key difference from the fitting in [21] was that instead of modeling the bolus input function by a trapezoid, this study used the measured Pyr time-curve directly as the bolus input into the model. As the measured Pyr time-curve includes the T_1 decay information, it obviates the need to separately fit for Pyr T_1 , and so apparent $T_{1\text{Pyr}}$ was fixed to 1 s. Good fits were obtained with apparent $T_{1\text{lac}} = 12 \text{ s}$ and $T_{1\text{ala}} = 15 \text{ s}$ for liver and kidney, and apparent $T_{1\text{lac}} = T_{1\text{ala}} = 20 \text{ s}$ for vasculature. These values were chosen empirically and kept the same for all animals.

The increase in apparent liver k_{pl} after infusion of ethanol (post-ethanol/pre-ethanol) observed here was $1.33 \pm 0.11 \text{ SEM}$ compared to the approximately 2-fold ($2.0 \pm 0.5 \text{ SEM}$) increase reported in [21]. This smaller effect may be due to the use of 80-mM Pyr instead of 100-mM Pyr as in that study, whereby the LDH enzyme activity may not be completely saturated. Though the total dose was comparable in both studies ($240 \mu\text{mol}$ vs. $250 \mu\text{mol}$), the lower concentration Pyr solution may be just at the boundary between the linear and saturated regimes of LDH enzyme activity, especially given the time-varying Pyr concentration during the $\sim 15 \text{ s}$ bolus injection.

The analysis in [21] used separate control measurements with saline injection instead of ethanol injection to confirm that the increased lactate production in the liver was an effect of ethanol. Based on those results, additional saline control measurements were not repeated for this study. The alanine kinetics as well as the lactate kinetics in kidney and in vasculature serve as self-control within each animal.

Conclusion

This work demonstrates fast volumetric imaging of hyperpolarized ^{13}C Pyr and its metabolic products *in vivo* using spectrally under-sampled spiral MRSI. The method was applied to measure the changes in Pyr-to-Lac conversion after infusion of ethanol. Spiral MRSI is an efficient acquisition method, achieving $80\times 80\times 60\text{ mm}^3$ volume coverage with a nominal 5-mm isotropic resolution in 4.5 s on a clinical 3T scanner. This implementation using clinical gradients makes the method suitable for translation to human use, e.g., the imaging parameters listed above would be suitable for prostate cancer imaging which is the target for the first clinical trial currently underway [23]. Combining this method with parallel imaging techniques would allow greater volume coverage, higher resolution, or even faster acquisition times, enabling other applications. Alternatively, a high performance insert gradient [11] can be used to further speed up the acquisition for small animal imaging, achieving a $2.7\times 2.7\times 5\text{-mm}^3$ resolution for the same $16\times 16\times 12$ matrix size in 1.5 s. The short acquisition times enable a high temporal resolution for monitoring dynamic processes. The dynamic 3D acquisition allows the analysis of spectra from organ-specific ROIs, thus providing improved rate-constant estimates and permitting quantitative evaluation of metabolic kinetics.

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Abbreviations used

MRSI	magnetic resonance spectroscopic imaging
FOV	field of view
ROI	region of interest
LDH	lactate dehydrogenase
NAD+	nicotinamide adenine dinucleotide
SPGR	spoiled gradient echo
BAL	blood alcohol level
SNR	signal-to-noise ratio

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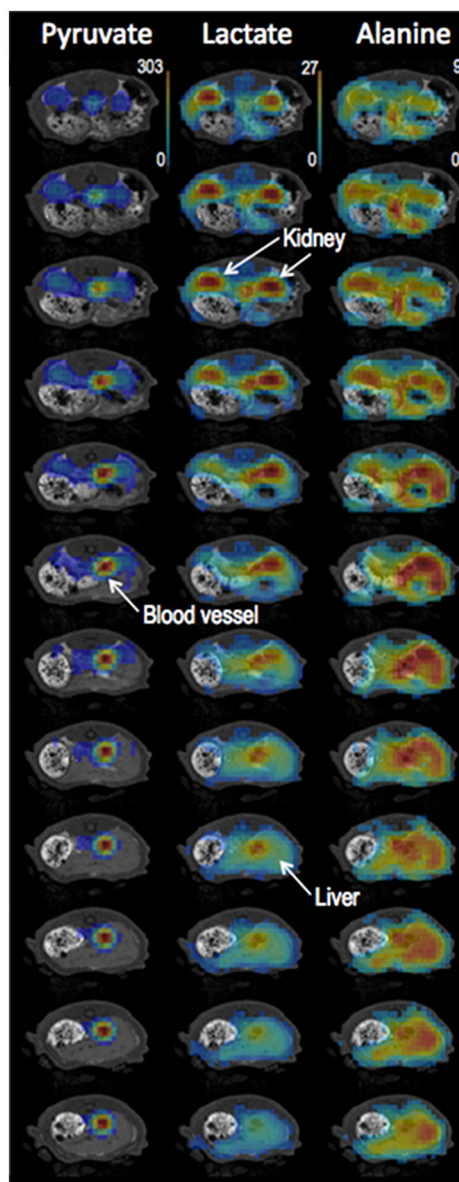


Figure 1. Representative post-ethanol ^{13}C metabolic maps of pyruvate, lactate and alanine superimposed on corresponding ^1H MRI images from the 3D volume covering kidneys and liver. The ^{13}C images are averaged over all 16 time-points acquired. The central 12 of the 24 reconstructed slices are shown here. The thresholds for the display of ^{13}C maps were 9% of the maximum intensity for pyruvate, and 20% for lactate and alanine.

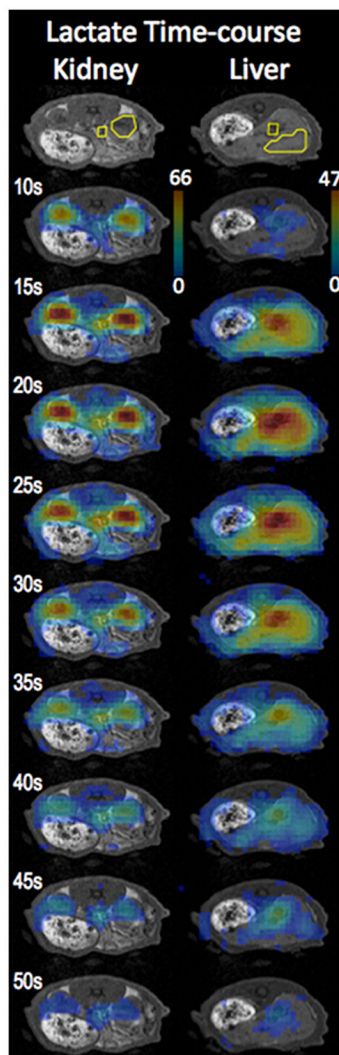


Figure 2.

Time series of lactate maps from kidney and liver slices illustrate the metabolite dynamics and distribution. The labeled time indicates time from beginning of the pyruvate injection. The threshold for the display of ^{13}C images was 15% of the maximum intensity. The ROIs used to obtain the metabolite time-courses are also shown.

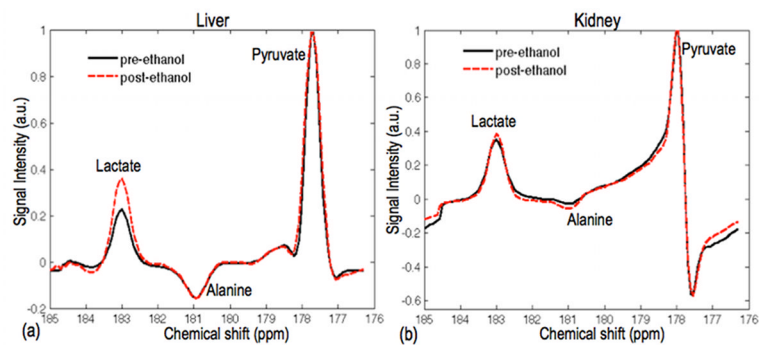


Figure 3. Representative time-averaged ^{13}C lactate spectra for liver (a) and kidney (b). The liver spectra show higher lactate signal post-ethanol compared to pre-ethanol while the kidney spectra showed similar lactate signal pre- and post-ethanol.

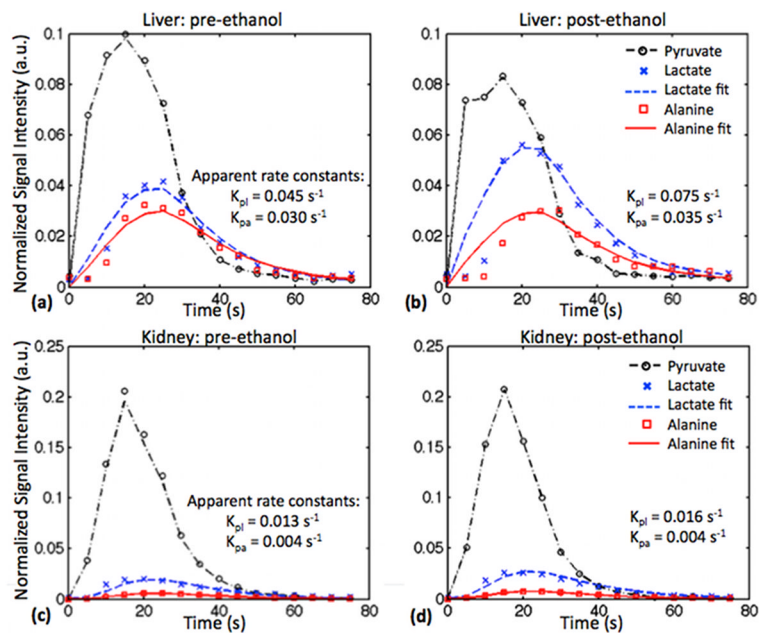
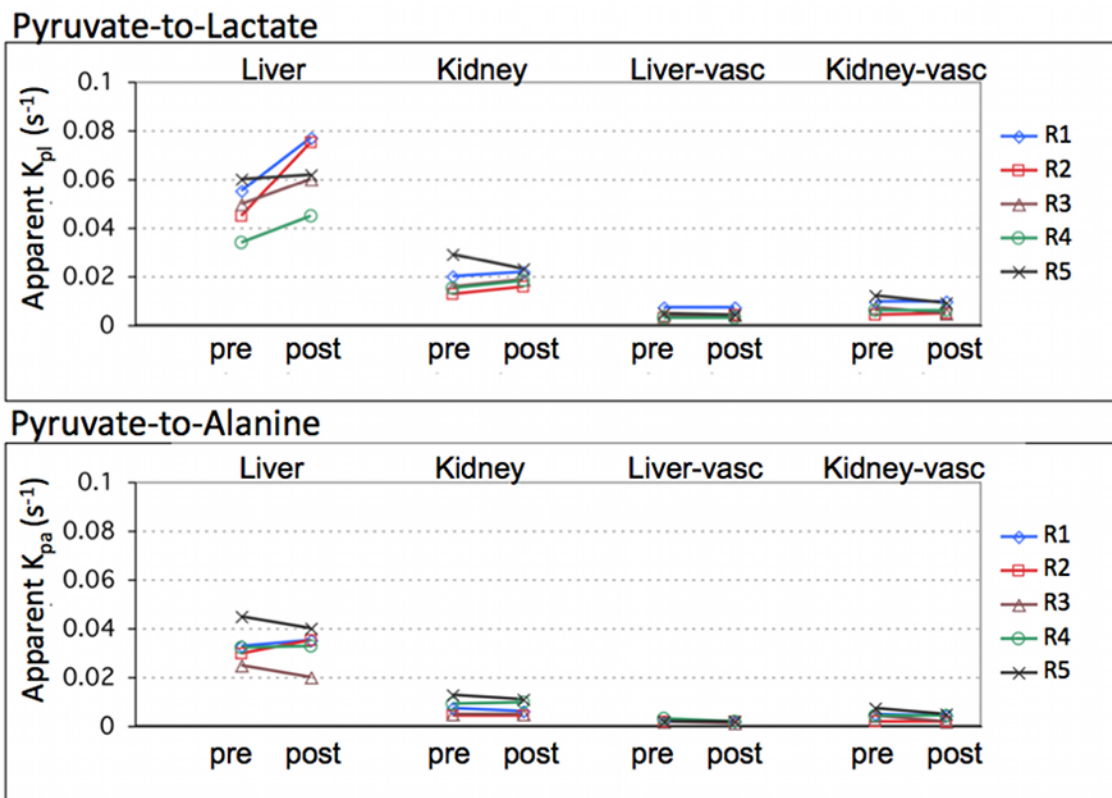


Figure 4.

Representative time course data, curve fits, and calculated apparent rate constants, from the liver acquired pre-ethanol (a) and 45 min post-ethanol (b), and from the kidney pre-ethanol (c) and post-ethanol (d). Higher lactate was measured in the liver after ethanol administration, whereas there was little change in kidney. The conversion of pyruvate to alanine was similar before and after ethanol in both organs.

**Figure 5.**

Estimated apparent k_{pl} and k_{pa} rate constants in liver, kidney, and vasculature in the respective slices. Conversion of labeled pyruvate to lactate increased in the liver after ethanol administration as seen by the increase in k_{pl} values. There was no significant change in k_{pl} in kidney or in vasculature or in k_{pa} .

Average apparent k_{pl} and k_{pa} rate constants with corresponding standard deviations in liver, kidney and vasculature ROIs from 5 animals. Conversion of labeled pyruvate to lactate increased in the liver after ethanol administration as seen by the increase in k_{pl} . There was no significant change in k_{pl} for kidney or vasculature, or in k_{pa} in any of the tissues. Though the inter-subject variability leads to relatively high standard deviations, the intra-subject comparison between pre- and post-ethanol in Fig. 5 clearly shows the change in rate constants.

Table 1

	Liver k_{pl} (s^{-1})	Liver k_{pa} (s^{-1})	Kidney k_{pl} (s^{-1})	Kidney k_{pa} (s^{-1})	Liver vasc. k_{pl} (s^{-1})	Liver vasc. k_{pa} (s^{-1})	Kidney vasc. k_{pl} (s^{-1})	Kidney vasc. k_{pa} (s^{-1})
Pre-ethanol	0.049 ± 0.009	0.033 ± 0.007	0.018 ± 0.004	0.008 ± 0.003	0.004 ± 0.002	0.002 ± 0.001	0.008 ± 0.003	0.004 ± 0.002
Post-ethanol	0.064 ± 0.012	0.032 ± 0.007	0.019 ± 0.003	0.007 ± 0.003	0.004 ± 0.002	0.002 ± 0.001	0.007 ± 0.002	0.003 ± 0.001
Wilcoxon p-value	0.0431	0.8927	0.4982	0.5930	1.0000	0.1573	0.2850	0.1025